

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Docket No.: 1/1144  
Application of: Jung, Birgit et al ) Art Unit: To be assigned  
Serial No. : To be assigned ) Examiner: To be assigned  
Filed : August 31, 2001  
For : Method for identifying substances which positively influence  
inflammatory conditions of chronic inflammatory airway disease

Assistant Commissioner for Patents  
Washington, D.C. 20231

STATEMENT BY ATTORNEY UNDER 37 C.F.R. § 1.821(f)  
REGARDING SEQUENCE LISTING

Sir:

Attorney for Applicants affirms that the information recorded in computer readable form is identical to the written sequence listing.

Respectfully submitted,

*Susan K. Pocchiaro*

Susan K. Pocchiaro  
Attorney for Applicant(s)  
Reg. No. 45,016

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900 Ridgebury Road  
P.O. Box 368  
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Assistant Commissioner for Patents  
Washington, DC 20231

on August 31, 2001

*Susan K. Pocchiaro*  
By: Susan K. Pocchiaro  
Reg. No. 45,016

**SEQUENCE LISTING**

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FATIGUE / 100+50

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 260 265 270  
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 275 280 285  
 His Gly Gly Gln Tyr Arg Cys Tyr Ser Ala His Asn Leu Ser Ser Glu  
 290 295 300  
 Trp Ser Ala Pro Ser Asp Pro Leu Asp Ile Leu Ile Thr Gly Gln Phe  
 305 310 315 320  
 Tyr Asp Arg Pro Ser Leu Ser Val Gln Pro Val Pro Thr Val Ala Pro  
 325 330 335  
 Gly Lys Asn Val Thr Leu Leu Cys Gln Ser Arg Gly Gln Phe His Thr  
 340 345 350  
 Phe Leu Leu Thr Lys Glu Gly Ala Gly His Pro Pro Leu His Leu Arg  
 355 360 365  
 Ser Glu His Gln Ala Gln Gln Asn Gln Ala Glu Phe Arg Met Gly Pro  
 370 375 380

Val Thr Ser Ala His Val Gly Thr Tyr Arg Cys Tyr Ser Ser Leu Ser  
385 390 395 400

Ser Asn Pro Tyr Leu Leu Ser Leu Pro Ser Asp Pro Leu Glu Leu Val  
405 410 415

Val Ser Ala Ser Leu Gly Gln His Pro Gln Asp Tyr Thr Val Glu Asn  
420 425 430

Leu Ile Arg Met Gly Val Ala Gly Leu Val Leu Val Val Leu Gly Ile  
435 440 445

Leu Leu Phe Glu Ala Gln His Ser Gln Arg Ser Leu Gln Asp Ala Ala  
450 455 460

Gly Arg  
465

<210> 13  
<211> 63  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer

<400> 13  
ggccagtgaa ttgtaatacg actcactata gggaggcgtt tttttttttt tttttttttt 60  
ttt 63

<210> 14  
<211> 25  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer

<400> 14  
gtcgtaaga tgctaccgtt cagga 25

<210> 15  
<211> 48  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer

<400> 15  
ggggacaagt ttgtacaaaa aagcaggcta tggaaaccaa cttctcca 48

<210> 16  
<211> 53  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer

<400> 16  
ggggaccact ttgtacaaga aagctgggtt cacattgcct gtaactcagt ctc 53

<210> 17  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer

<400> 17  
agcccatagc agatggcaac 20

NOTE: 208nt=60

<210> 18  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer

<400> 18  
tgtactttca actttgcata ctgg 24

<210> 19  
<211> 28  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer

<400> 19  
aagccaatga caaacggat aatccctc 28

<210> 20  
<211> 2051  
<212> DNA  
<213> Homo sapiens

<400> 20  
cgccactttg ctggaggcatt cactaggcga ggcgcctccat cggaacctact agccgcactc 60  
atgaatcgcc accatctgca ggatcactt ctggaaatag acaaaaagaa ctgctgtgt 120  
ttccgagatc attcatatgc caagggtttt ccggccgtgt tggggctgga gtttatcttt 180

gggcttctgg gcaatggcc tgcctgtgg attttctgtt tccaccaa gtctggaaa 240  
 tccacccgga ttttccgtt caactccgaa stagtgact ttctactgt catctgcgtc 300  
 cogttctgtg tggactacta tggtcgccgt tcagactgga actttggga cattccctgc 360  
 cggtctgtc ttctcatgtt tgccatgaa cgcaggcca gcatactt ctcacagggt 420  
 gtggcggtg acagattttt ccgggtggto catccccacc acgocctgaa caagacttcc 480  
 aattggacac cagccatcat ctcttgcctt ctgtggcca tcacttgtt cctaacaagt 540  
 cacccctcgtg agagaagggt gctgtatccg aatggccctg caaaatgtgtg catcagcttc 600  
 agcatctggcc atacccctccg tgggccacgaa gctatgttc tcttgaggtt ctcctgccc 660  
 ctgggcataca tcctgttctg ctcagccaga attatctggaa gcctgcggca gagacaatg 720  
 gaccggcgtc ccaagatccaa gagaggccatc accttcatca ttgtgtggc catcgcttt 780  
 gtcatctgtc ttcttcccg ctgtgtgtg cggatccgaa ttcttcgtt cttgcacact 840  
 tcgggcacgc aagaattgtga agtgtacccg ctgtggacc ttggcttctt tatcacttc 900  
 agcttcacccat acatggacag catgtggac ccctgtgtt actacttctc cagccatcc 960  
 ttttccaaact ttcttcctcaat ttgtatcaac cgctgccttc agaggaaatg gagagggttag 1020  
 ccagataaa accgcagccgac gacgcgtccg ctcaacagggg accccaaacaa aaccaggaggc 1080  
 gtcctccaggcg ctgtatggc caactccgtt gageccatggaa gccccttta tctggggccca 1140  
 acctcaataa accatcccaaa gaaaggacat ttgtaccaag aaccaggatc tctggagaaa 1200  
 cagttgggctt gtgtacccgt gtaatgtcaat ggacttcggc ctaagggttcc ttggaaacttc 1260  
 cagattcaga gaatctgatt tagggaaact gtggcagatg agtgggagac ttgttgtcaag 1320  
 gtgtgaccac aggaatcttccg gaggaaacaga gatgtaaatggc tcttaggcattc tgaacttc 1380  
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 ggggggggctt cagctccctcg ggtatatatc agccctgtt tgactcttag caggataaag 1560  
 gagactgtcgg attggggggggc attgtgttc tcttgaggaa agccaggccaa tcattaaaca 1620  
 agccatagggttccatctgttcc tccgtggacc aattatctt tcagacaaaggc tttagaaaaaa 1680  
 tggactctggaaatggggacttccatctgttcc tttatggaaa gaaatggca 1740  
 aggggtaggttcccaaaaggc gactgtggca aacagtgttta ttatggggaaa gaaatggca 1800  
 ttgtgttccatctgttccatctgttcc tccatcttcc tttatggggaaa gaaatcttaa 1860  
 ggttgaggccat taaacccggc ttccatgttcc aacagtgttcc tccacccgtt ttgttttacc 1920  
 attaaaaaaagggaaatccgtcc tcccccacccggttagggggggc ttccatctgttcc tccatctgttcc 1980  
 ctccatctgttccatctgttcc tccatctgttcc tccatctgttcc tccatctgttcc 2040  
 caaaaaaaaaaa a 2051

100  
 150  
 200  
 250  
 300  
 350  
 400  
 450  
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 550  
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<210> 21  
 <211> 397  
 <212> PRT  
 <213> Homo sapiens

<400> 21  
 Met Asn Arg His His Leu Gln Asp His Phe Leu Glu Ile Asp Lys Lys  
 1 5 10 15

Asn Cys Cys Val Phe Arg Asp Asp Phe Ile Ala Lys Val Leu Pro Pro  
 20 25 30

Val Leu Gly Leu Glu Phe Ile Phe Gly Leu Leu Gly Asn Gly Leu Ala  
 35 40 45

Leu Trp Ile Phe Cys Phe His Leu Lys Ser Trp Lys Ser Ser Arg Ile  
 50 55 60

Phe Leu Phe Asn Leu Ala Val Ala Asp Phe Leu Leu Ile Ile Cys Leu  
 65 70 75 80

Pro Phe Val Met Asp Tyr Tyr Val Arg Arg Ser Asp Asp Trp Asn Phe Gly  
 85 90 95

Asp Ile Pro Cys Arg Leu Val Leu Phe Met Phe Ala Met Asn Arg Gln  
 100 105 110  
 Gly Ser Ile Ile Phe Leu Thr Val Val Ala Val Asp Arg Tyr Phe Arg  
 115 120 125  
 Val Val His Pro His His Ala Leu Asn Lys Ile Ser Asn Trp Thr Ala  
 130 135 140  
 Ala Ile Ile Ser Cys Leu Leu Trp Gly Ile Thr Val Gly Leu Thr Val  
 145 150 155 160  
 His Leu Leu Lys Lys Leu Leu Ile Gln Asn Gly Pro Ala Asn Val  
 165 170 175  
 Cys Ile Ser Phe Ser Ile Cys His Thr Phe Arg Trp His Glu Ala Met  
 180 185 190  
 Phe Leu Leu Glu Phe Leu Leu Pro Leu Gly Ile Ile Leu Phe Cys Ser  
 195 200 205  
 Ala Arg Ile Ile Trp Ser Leu Arg Gln Arg Gln Met Asp Arg His Ala  
 210 215 220  
 Lys Ile Lys Arg Ala Ile Thr Phe Ile Met Val Val Ala Ile Val Phe  
 225 230 235 240  
 Val Ile Cys Phe Leu Pro Ser Val Val Arg Ile Arg Ile Phe Trp  
 245 250 255  
 Leu Leu His Thr Ser Gly Thr Gln Asn Cys Glu Val Tyr Arg Ser Val  
 260 265 270  
 Asp Leu Ala Phe Phe Ile Thr Leu Ser Phe Thr Tyr Met Asn Ser Met  
 275 280 285  
 Leu Asp Pro Val Val Tyr Tyr Phe Ser Ser Pro Ser Phe Pro Asn Phe  
 290 295 300  
 Phe Ser Thr Leu Ile Asn Arg Cys Leu Gln Arg Lys Met Thr Gly Glu  
 305 310 315 320  
 Pro Asp Asn Asn Arg Ser Thr Ser Val Glu Leu Thr Gly Asp Pro Asn  
 325 330 335  
 Lys Thr Arg Gly Ala Pro Glu Ala Leu Met Ala Asn Ser Gly Glu Pro  
 340 345 350  
 Trp Ser Pro Ser Tyr Leu Gly Pro Thr Ser Asn Asn His Ser Lys Lys  
 355 360 365  
 Gly His Cys His Gln Glu Pro Ala Ser Leu Glu Lys Gln Leu Gly Cys  
 370 375 380  
 Cys Ile Glu  
 385

<210> 22  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer

<400> 22  
agcccatagc agatggcaac 20

<210> 23  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer

<400> 23  
tgtactttca actttgcata ctgg 24

<210> 24  
<211> 28  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer

<400> 24  
aagccaatga caaacggat aatccctc 28